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Research Article

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Regulation of myeloproliferation and M2 macrophage programming in mice by Lyn/Hck, SHIP, and Stat5

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The proliferation and differentiation of hematopoietic stem cells (HSCs) is finely regulated by extrinsic and intrinsic factors via various signaling pathways. Here we have shown that, similar to mice deficient in the lipid phosphatase SHIP, loss of 2 Src family kinases, Lyn and Hck, profoundly affects HSC differentiation, producing hematopoietic progenitors with increased proliferation, reduced apoptosis, growth factor-independent survival, and skewed differentiation toward M2 macrophages. This phenotype culminates in a Stat5-dependent myeloproliferative disease that is accompanied by M2 macrophage infiltration of the lung. Expression of a membrane-bound form of SHIP in HSCs lacking both Lyn and Hck restored normal hematopoiesis and prevented myeloproliferation. In vitro and in vivo studies suggested the involvement of autocrine and/or paracrine production of IL-3 and GM-CSF in the increased proliferation and myeloid differentiation of HSCs. Thus, this study has defined a myeloproliferative transformation-sensitive signaling pathway, composed of Lyn/Hck, SHIP, autocrine/paracrine cytokines, and Stat5, that regulates HSC differentiation and M2 macrophage programming.

Introduction

The Src family kinases (SFKs) mediate signals from a multitude of cell surface receptors to regulate a variety of cellular functions including proliferation, differentiation, and migration (1). Several SFK members are expressed in hematopoietic cells and linked to signaling by a number of cytokine receptors and receptor tyrosine kinases. Thus, SFKs play important roles in the development and function of hematopoietic cells: for example, *lyn*^{-/-} mice suffer from a deficit of mature B cells (2, 3). However, aged *lyn*^{-/-} mice display autoimmune disease and perturbations in myelopoiesis that ultimately lead to splenomegaly and macrophage tumors (4). Impaired hematopoiesis in *lyn*^{-/-} mice is explained by the essential role of Lyn in establishing immunoreceptor tyrosine-based inhibitory motif-dependent (ITIM-dependent) signaling. ITIM-containing inhibitory receptors phosphorylated by Lyn recruit protein tyrosine phosphatases, such as SH2-domain containing phosphatase-1 (SHP-1) and SHP-2 as well as SH2-domain containing 5'-inositol phosphatase (SHIP-1), to the plasma membrane (5). Perturbations in these signaling pathways result in an increased number of myeloid progenitors (6) and impaired B-lymphopoiesis in *lyn*^{-/-} mice. The SFKs Hck and Fgr also serve an inhibitory function in hematopoiesis and mature leukocyte function. Deficiency of Hck alone results in increased G-CSF induced proliferation of granulocytic precursors (6), while loss of both kinases results in

exaggerated chemokine-induced activation in neutrophils and dendritic cells (7). Despite these observations, roles of SFKs in HSC homeostasis are poorly defined.

SHIP is expressed primarily in hematopoietic cells including HSCs (8, 9). This enzyme hydrolyzes the 5' phosphate on phosphatidylinositol 3,4,5-trisphosphate and inositol 1,3,4,5-tetrakisphosphate, thus preventing membrane recruitment and activation of pleckstrin homology domain-containing kinases that serve as effectors of phosphatidylinositol-3'-kinase (PI3K) signaling. SHIP associates with various adaptor/scaffold proteins or inhibitory receptors following activation of hematopoietic cells. SHIP-deficient mice exhibit chronic hyperplasia of myeloid cells resulting in splenomegaly, lymphadenopathy, and myeloid infiltration of lung and other organs (10, 11). Neutrophils and bone marrow-derived (BM-derived) mast cells from *SHIP*^{-/-} mice are resistant to apoptosis. Consistent with these observations, HSC proliferation and numbers are increased in *SHIP*^{-/-} mice (12). Thus, SHIP limits the survival, activation, proliferation, and differentiation of hematopoietic cells.

Here we show that mice deficient in Lyn and Hck develop myeloproliferative disease (MPD) accompanied by severe lung inflammation with extensive macrophage infiltration, and that both the MPD and the lung inflammation originate from long-term HSCs (LT-HSCs). Our study has also revealed a signaling pathway for Lyn/Hck/SHIP-mediated inhibition of cytokines production that, via the Stat5 signaling, tightly controls the proliferation, apoptosis, and differentiation of HSCs and myeloid progenitors. Perturbations of this pathway lead to myeloproliferative transformation of LT-HSCs.

Results

Lyn/Hck-deficient mice suffer from severe lung inflammation. During our study of *lyn*^{-/-}, *hck*^{-/-}, and *lyn*^{-/-}*hck*^{-/-} (double-knockout [dko]) mice, we noticed that dko mice died prematurely. Dko mice were

Nonstandard abbreviations used: BAL, bronchoalveolar lavage; CML, chronic myelogenous leukemia; CMP, common myeloid progenitor; dko, double-knockout; DN, dominant-negative; G-CFU, granulocyte-CFU; GMP, granulocyte-macrophage progenitor; KSL, c-Kit⁺ Sca-1⁺ lineage; LT-HSC, long-term HSC; Mix-CFU, granulocyte/erythrocyte/monocyte/megakaryocyte-CFU; MPD, myeloproliferative disease; SFK, Src family kinase; SHIP, SH2-domain containing 5'-inositol phosphatase; SHP-1, SH2-domain containing phosphatase-1; ST-HSC, short-term HSC.

Conflict of interest: The authors have declared that no conflict of interest exists.

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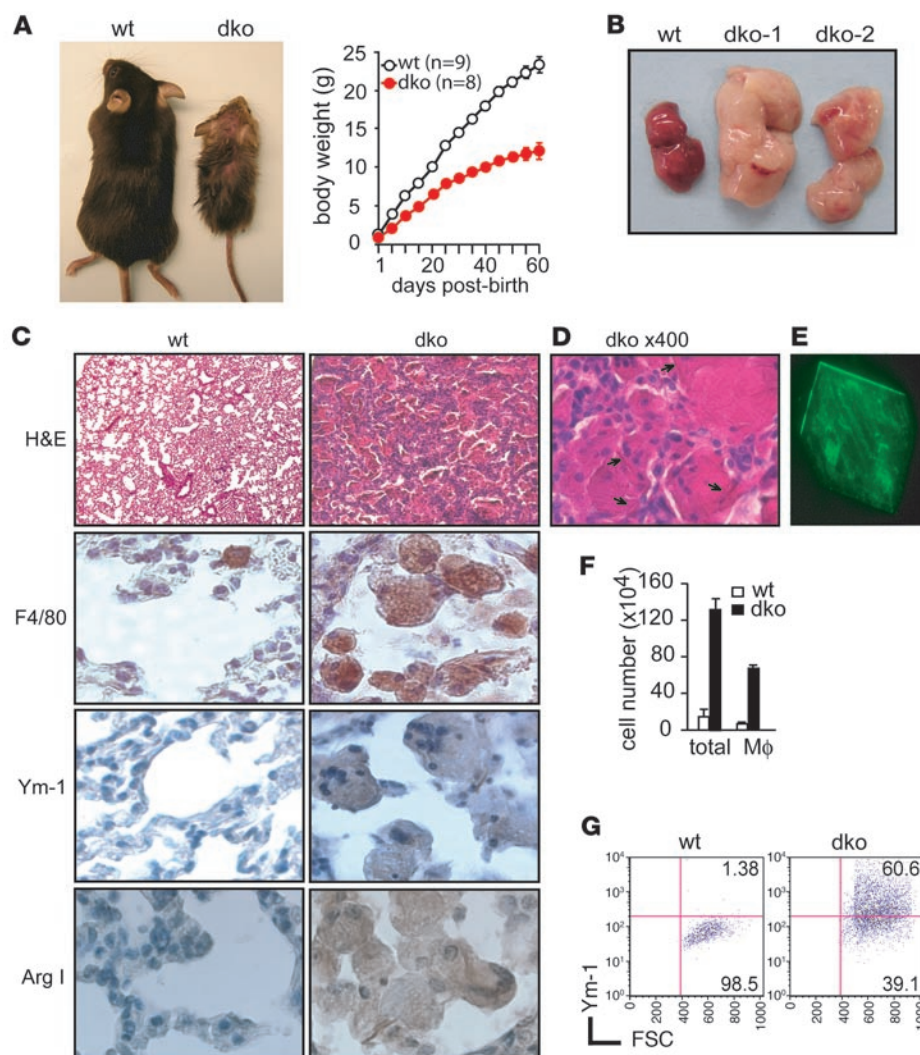


Figure 1

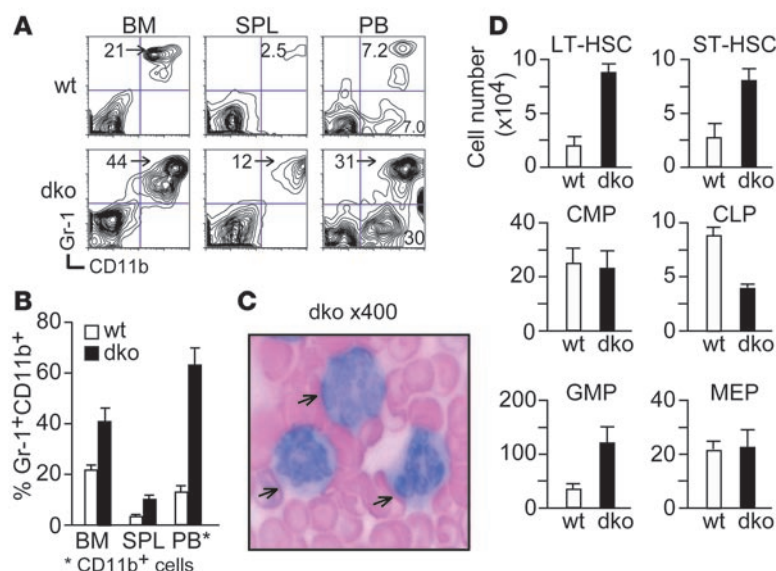
Severe lung inflammation characterized by extensive infiltration with M2-type macrophages in *Lyn/Hck*-deficient mice. (A) Growth of dko mice was slower than WT mice. The mice shown are 8 weeks in age. Data represent mean \pm SD throughout the figures. (B) Gross appearance showed that lungs are consolidated in dko mice. (C and D) Microscopic examination showed abundant M2 macrophages in alveoli of dko lungs. Original magnification, $\times 100$ (H&E); others, $\times 400$ (C). Arrows indicate crystals. (E) A crystal recovered from BAL from dko mice and illuminated by green light of 488 nm (original magnification, $\times 400$). (F) Increased numbers of macrophages ($M\phi$) were recovered in BAL fluids of dko mice. (G) Ym-1-positive cells were increased in BAL macrophages from dko mice.

born at normal Mendelian frequency (Supplemental Figure 1; supplemental material available online with this article; doi:10.1172/JCI34013DS1); however, within 2 weeks they became obviously sick, growth retarded, and most died within 2 months (Figure 1A). By contrast, *lyn*^{-/-} and *hck*^{-/-} mice remained healthy for the first 2 months of age (data not shown). Gross examination revealed that the lungs in dko mice were bilaterally consolidated (Figure 1B). Histologically, the lungs were fibrotic and profoundly infiltrated with inflammatory cells, the majority of which were F4/80⁺ macrophages (Figure 1, C and D, and Supplemental Figure 2). Compared with WT mice, total cell numbers in bronchoalveolar lavages (BALs) were increased by 10-fold in dko mice (Figure 1F). Fifty-five percent of the cells were macrophages, and most other cells (32%) were CD11b⁺Gr-1⁺F4/80⁺ monocytes (data not shown). These macrophages expressed Ym-1 and arginase I (Figure 1, C and G), suggesting an M2-skewed phenotype (13, 14). Similar to those in *SHIP*^{-/-} mice (10, 11, 15), eosinophilic hexagonal crystals were abundantly found in BAL fluids recovered from dko, but not WT, mice (Figure 1E). Given the extensive nature of lung inflammation, this could be a cause of premature deaths in dko mice.

Dko mice develop myeloproliferative disease. In addition to the macrophage/monocyte pulmonary inflammation, dko mice displayed

splenomegaly (Supplemental Figure 3, A and B). A dramatic increase of granulocytes in BM, spleen, and peripheral blood from dko mice was revealed by flow cytometric analysis (Figure 2, A and B) and differential blood cell counting (Supplemental Figure 3D). Leukocyte numbers in peripheral blood were increased by 3-fold ($21,750 \pm 3,780/\mu\text{l}$ in dko mice versus $6,850 \pm 540/\mu\text{l}$ in WT mice; $P < 0.001$; $n = 5$) in 2-month-old mice. Most of the increased leukocytes were mature granulocytes (Figure 2C). Monocytes (CD11b⁺Gr-1⁺) in peripheral blood were also increased (Figure 2A and Supplemental Figure 3D). In addition to pulmonary infiltration, dko mice also showed evidence of hepatic infiltration by myeloid cells (Supplemental Figure 3C). These data collectively indicate that all dko mice analyzed ($n = 12$) exhibit a set of clinical signs within 2 months of age that fulfill the criteria for the MPD (16). No abnormalities in myelopoiesis were observed with either *lyn*^{-/-} or *hck*^{-/-} mice for the first 2 months after birth, whereas old (>6 months of age) *lyn*^{-/-}, but not *hck*^{-/-}, mice showed a mild MPD phenotype (ref. 17 and data not shown).

Dko mice have increased HSCs and myeloid progenitors. Given their MPD phenotype, we investigated the hematopoietic process in dko mice in more detail. Flow cytometric analysis revealed that the absolute number of c-Kit⁺ Sca-1⁺ lineage⁻ (KSL) cells, which are

**Figure 2**

The MPD phenotype in *Lyn/Hck*-deficient mice. (A and B) Increased myeloid cells (Gr-1⁺CD11b⁺ or CD11b⁺) were found in BM, spleen (SPL), and peripheral blood (PB) from dko mice, as quantified by flow cytometry. (C) Shown is a Wright-Giemsa-stained blood smear of dko mouse (original magnification, ×400). Arrows indicate neutrophils. (D) Accumulated flow cytometric data of BM cells are shown (*n* = 12). Numbers of specific cell types were calculated by multiplying their percentages (obtained by flow cytometry; see Supplemental Figure 4) by total cell numbers. Dko mice have increased numbers of HSCs and myeloid progenitors but decreased numbers of lymphoid progenitors (see Supplemental Figures 4 and 5). MEP, megakaryocyte-erythroid progenitor.

enriched for HSCs, in BM from dko mice is increased by 4.1-fold compared with WT mice (Figure 2D and Supplemental Figure 4). Further, dko mice had 4.2-fold more KSL CD34⁺ cells (enriched for long-term HSCs [LT-HSCs]; ref. 18) and 3.2-fold more KSL CD34⁺ cells (enriched for short-term HSCs [ST-HSCs]). Granulocyte-macrophage progenitors (GMPs) were also increased by 3-fold. However, both common myeloid progenitors (CMPs) and megakaryocyte-erythroid progenitors (MEPs) were comparable in WT and dko mice. By contrast, the number of common lymphoid progenitors was decreased by 2-fold (Figure 2D), consistent with reduced T and B cells in dko mice (Supplemental Figure 5). Compared with WT cells, BM cells from dko mice formed increased numbers of myeloid colonies (i.e., granulocyte-CFU [G-CFU], M-CFU, GM-CFU, and granulocyte/erythrocyte/monocyte/megakaryocyte-CFU [Mix-CFU]) in methylcellulose medium containing IL-3, IL-6, stem cell factor (SCF), and erythropoietin, but reduced numbers of pre-B cell colonies in medium containing IL-7 (Supplemental Figure 4, B and C). Overall, *lyn/hck* dko mice had increased HSCs and GMPs, similar to other mouse MPD models (19–21), suggesting that the MPDs in these mice may be caused by similarly perturbed hematopoiesis (Supplemental Figure 4D).

MPD and lung disease originate from LT-HSCs in dko mice. A cardinal feature of the mouse MPD is the transplantability of disease (19). Therefore, we performed a series of transplantation experiments (Table 1). First, BM cells from WT or dko mice (Ly5.2) were i.v. injected into lethally

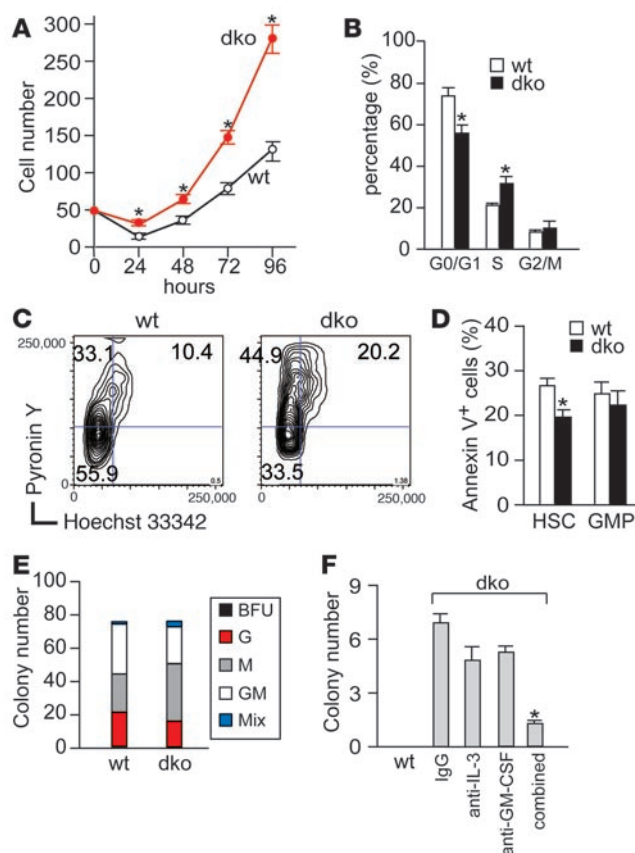
irradiated C57BL/6-Ly5.1 mice. All the mice that had received dko BM cells developed MPD and lung disease and died within 2 months. Histological examination confirmed that the lung disease in recipients engrafted with dko BM cells was indistinguishable from that in donor dko mice (Supplemental Figure 6), indicating

Table 1

MPD and lung inflammation are transplantable with LT-HSCs of *Lyn/Hck*-deficient mice

Donor cells	Engraftment ^A		% Gr-1 ⁺ CD11b ⁺	Myeloid hyperplasia/ lung infiltration per recipient (no.)
	3 wk	9 wk		
BM cells				
WT	nd	>99	18.5 ± 5.2	0/8
dko	nd	>99	71.0 ± 11.0	12/12
LT-HSC				
WT	54.0 ± 11	58.0 ± 15	22.0 ± 4.3	0/6
dko	46.7 ± 4.3	49.1 ± 5.6	78.4 ± 3.3	8/8
dko+vec	33.0 ± 13	38.0 ± 14	68.0 ± 8.7 ^B	6/6
dko+DN5	10.0 ± 0.8	9.0 ± 1.8	21.4 ± 6.2 ^B	0/8
ST-HSC				
WT	51.2 ± 3.2	11.4 ± 4.5	5.9 ± 2.6	0/4
dko	41.0 ± 9.4	9.0 ± 5.1	4.1 ± 1.5	0/4
CMP				
WT	3.2 ± 1.6	0	0	0/4
dko	3.4 ± 0.6	0	0	0/4
GMP				
WT	8.3 ± 3.5	0	0	0/4
dko	0.8 ± 0.4	0	0	0/4

The MPD and lung phenotypes were transplantable with BM or KSL CD34⁺ cells (LT-HSCs). These phenotypes were prevented by expression of DN Stat5 in LT-HSCs. BM cells (2 × 10⁶), LT-HSCs (1 × 10⁴), ST-HSCs (2 × 10⁴), CMPs (2 × 10⁴), GMPs (2 × 10⁴), or transduced LT-HSCs were i.v. transferred into lethally irradiated Ly5.1 mice. LT-HSCs and other fractionated cells were cotransferred with 2 × 10⁵ Sca-1⁺ BM cells of Ly5.1 mice. Peripheral blood cells were analyzed 3 and 9 weeks after transplantation. The fourth column shows percent Gr-1⁺CD11b⁺ cells among the donor-derived cells at the ninth week as determined by staining for Ly5.2. Dying mice were sacrificed, and the MPD phenotype was confirmed by flow cytometry of BM and spleen cells. Lung inflammation was detected by histological analysis. All the mice that developed MPD also developed lung infiltration with macrophage infiltrates. Transplantation results were combined from 2 independent experiments. ^A%CD45.2⁺ cells. ^BCD11b⁺ cells.

**Figure 3**

HSCs from dko mice exhibit increased proliferation, decreased apoptosis, and skewed differentiation into macrophages. (A) KSL CD34⁺ cells derived from WT and dko mice were sorted into a 96-well plate containing SCF, Flt3 ligand, and IL-11, and cultured for 4 days. * $P < 0.05$ versus WT values by Student's t test. (B and C) Cell cycle analysis of KSL cells was performed by flow cytometry of cells stained with propidium iodide (B) or with Hoechst 33342 and Pyronin Y (C). Lower left, upper left, and upper right quadrants in panel C contain cells in G₀, G₁, and S/G₂/M phases, respectively. G₀, WT 55.1% \pm 0.8% versus dko 34.2% \pm 0.7%, $P = 0.002$; G₁, 31.8% \pm 1.4% versus 45.0% \pm 0.1%, $P = 0.01$; S/G₂/M, 10.1% \pm 0.3% versus 20.9% \pm 0.7%, $P = 0.004$. (D) Flow cytometric analysis indicated reduced annexin V⁺ apoptotic cells in KSL (HSC), but not in GMPs, derived from BM of dko mice. (E) Colony-forming assays showed skewed differentiation of KSL cells (100 cells seeded) of dko mice into macrophages in the presence of IL-3, IL-6, SCF, and erythropoietin. (F) Dko KSL cells (50 cells/well seeded), unlike WT cells, formed colonies in methylcellulose without growth factors, which was inhibited by anti-IL-3 and anti-GM-CSF antibodies. BFU, BFU-E; G, G-CFU; M, M-CFU; GM, GM-CFU; Mix, Mix-CFU.

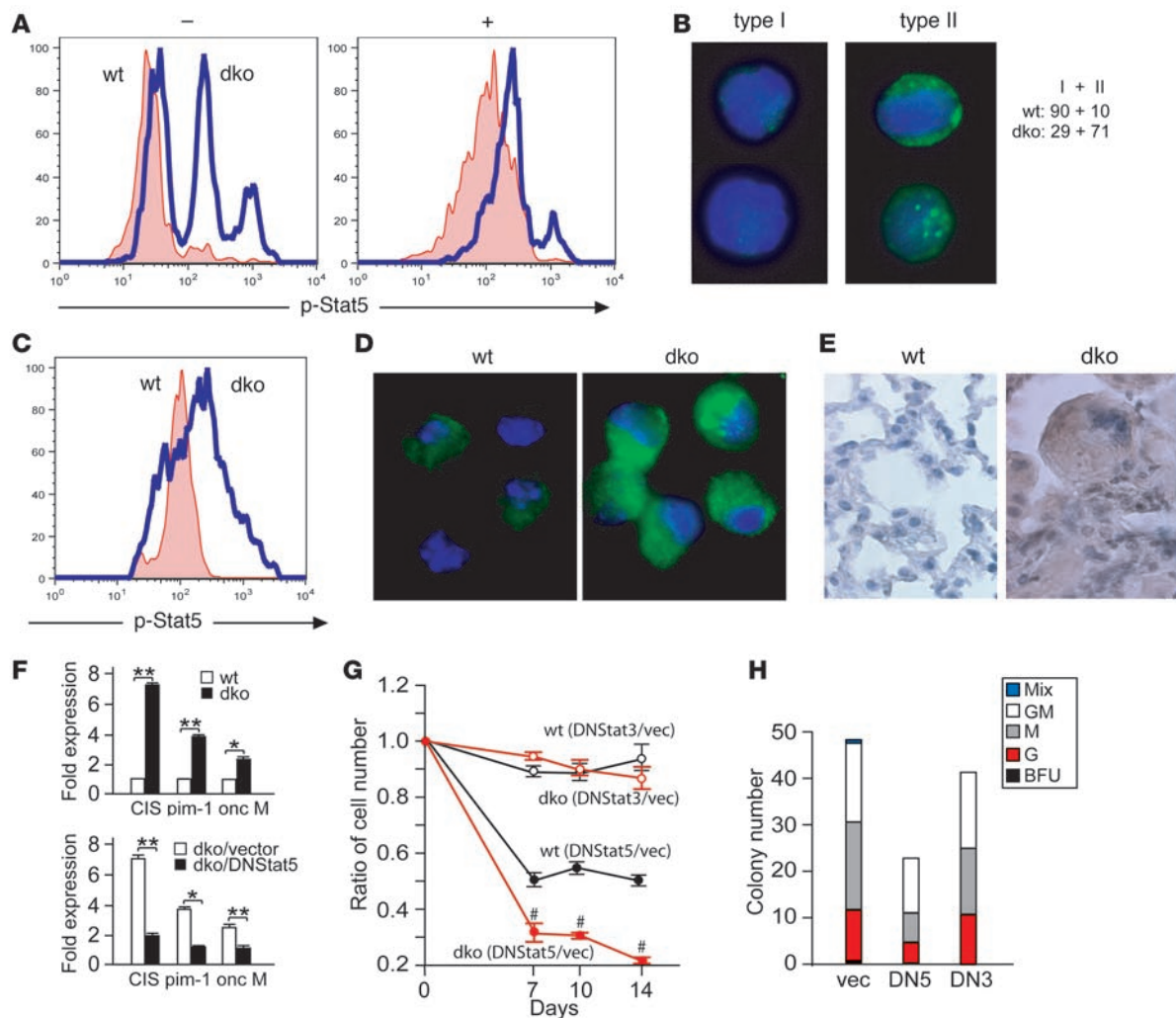
that both the MPD and lung disease are BM cell-autonomous. By contrast, none of the mice that had received WT BM cells developed MPD or lung disease. Next, purified subpopulations of BM cells from dko or WT mice were injected into lethally irradiated C57BL/6-Ly5.1 mice to identify the precise BM compartment that is responsible for the development of MPD and lung disease in dko mice. All the mice that had received dko, but not WT, LT-HSCs developed both MPD and lung disease within 9 weeks (Table 1). By contrast, the mice that had received ST-HSCs, CMPs, or GMPs derived from dko or WT mice failed to develop either disease, although there were no significant differences in engraftment of LT-HSC, ST-HSC, or CMP between WT and dko recipients 3 weeks after transplant (Table 1). The engraftment of GMP derived from dko mice was even lower than that of WT GMP. These results strongly suggest that the MPD and lung inflammation in dko mice originates from LT-HSCs, although we cannot rule out the possibility that other hematopoietic populations of dko mice might have the ability to cause the diseases. For example, transplantation with larger numbers of ST-HSCs or myeloid progenitors of dko mice might be able to cause the diseases in recipient mice.

HSCs from dko mice proliferate and differentiate into macrophages more vigorously and apoptose less abundantly than WT HSCs. The increase of HSCs in dko mice could be due to an increased proliferation, decreased apoptosis, altered differentiation, or a combination of these factors. In fact, both KSL cells and KSL CD34⁺ cells from dko mice grew faster under various in vitro culture conditions than WT cells (Figure 3A and data not shown). Consistent with this, cell cycle analysis showed increased proportions of dko KSL cells in the G₁ and S phases and reduced proportions in the G₀

phase (Figure 3, B and C). Moreover, fewer numbers of dko KSL cells stained with annexin V, indicating that dko HSCs are more resistant to apoptosis (Figure 3D). Finally, colony-forming assays showed that dko KSL cells generated more monocyte/macrophage (M-CFU) colonies than WT cells, indicating that dko HSCs have an increased predisposition to differentiate into monocytes and macrophages (Figure 3E). Moreover, a small population (~14%) of KSL cells formed GM-CFU colonies spontaneously in the absence of growth factors (Figure 3F), indicating that these progenitors contained some transformed cells. Taken together, the proliferation, survival, differentiation, and myeloproliferative transformation of HSCs are tightly regulated by Lyn and Hck.

Increased Stat5 activation is essential for increased proliferation and granulocyte/macrophage differentiation of HSCs in dko mice. To dissect the molecular mechanism for the increased proliferation, survival, and myeloid differentiation of dko HSCs, we examined several signaling pathways, including MAPKs, Akt, and Jak-Stat, which work as signal mediators for proliferation and survival of HSCs (22, 23). Although there is no significant difference in phosphorylation of ERK1/2, p38, and Stat3 (Supplemental Figure 7), phosphorylation of Akt, a serine/threonine kinase critical for cell survival (reviewed in ref. 24), was modestly increased in dko KSL cells compared with WT cells, consistent with the modestly increased survival of the former cells. Importantly, phosphorylation levels of Stat5 at Tyr694 were dramatically increased in unstimulated and cytokine-stimulated dko KSL cells (Figure 4A). Flow cytometry and imaging analysis demonstrated that KSL cells with nuclear localization of phosphorylated Stat5 (type II cells in Figure 4B) were more frequently found in dko mice. Similar results with Stat5 phosphorylation were also obtained with BAL macrophages from dko mice (Figure 4, C–E). mRNA and protein levels of Stat5 (both Stat5A and Stat5B) were comparable in dko and WT KSL cells (data not shown). Consistent with Stat5 activation, mRNA levels of the well-known Stat5 targets, e.g., cytokine-inducible SH2-containing protein (CIS) (25), pim-1 (26) (a serine/threonine kinase that can promote cytokine-independent survival; ref. 27), and oncostatin M (28), were all increased in dko cells (Figure 4F, upper panel). These results indicate that Stat5 is constitutively activated in dko HSCs.

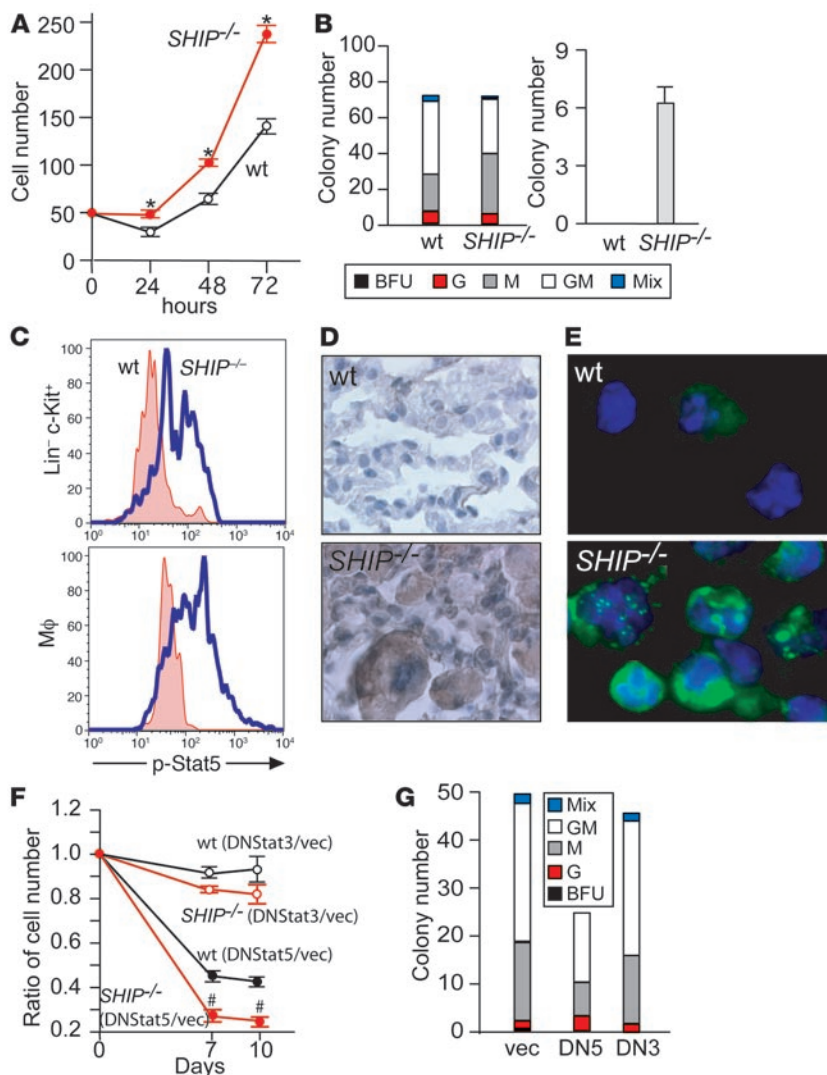
To directly test the functional relevance of the increased Stat5 activity, we introduced dominant-negative (DN) Stat5 into WT

**Figure 4**

Stat5 activation is required for the increased proliferation and macrophage differentiation of KSL cells from Lyn/Hck-deficient mice. (**A** and **B**) Stat5 in dko KSL cells was constitutively activated (–), and its activity was further increased upon stimulation with a cytokine cocktail of IL-3, SCF, and Flt3 ligand (+), as shown by flow cytometric analysis (**A**) and confocal microscopy (**B**) using anti-p-Stat5 (Tyr694) antibody. Nuclei were stained with DAPI (blue). Original magnification, $\times 400$. One hundred cells were scored for either the absence (type I) or presence (type II) of anti-p-Stat5 staining and those numbers are shown to the right of the panel. (**C–E**) Stat5 in dko lung macrophages was constitutively phosphorylated, as shown by flow cytometric analysis (**C**), confocal microscopy (**D**), and immunohistochemistry (**E**). Original magnification, $\times 400$ (**D** and **E**). (**F**) Real-time PCR was performed for mRNA expression of Stat5 target genes, cytokine-inducible SH2-containing protein (*CIS*), *pim-1*, and *oncostatin M* (*onc M*). Expression was normalized against L32 RNA and set at 1 in WT cells. * $P < 0.05$; ** $P < 0.01$. (**G**) Dko and WT KSL cells were transduced with a bicistronic retroviral vector encoding DN Stat5, DN Stat3, or empty vector, together with GFP, and cultured in medium containing SCF, Flt3 ligand, and IL-11. Ratios of cell numbers normalized against those of empty vector-transduced cells are plotted. For example, “WT (DNStat5/vec)” indicates the ratio of the numbers of WT cells transduced by DN Stat5 to those of WT cells transduced by empty vector. # $P < 0.05$ versus WT (DNStat5/vec). (**H**) Transduced KSL cells (150 cells seeded) were cultured in methylcellulose medium containing IL-3, IL-6, SCF, and erythropoietin. BFU, BFU-E; G, G-CFU; M, M-CFU; GM, GM-CFU; Mix, Mix-CFU.

and dko KSL cells, using a bicistronic retrovirus encoding DN Stat5 (DN Stat3 or empty vector as controls) and green fluorescent protein (GFP). As expected, DN Stat5 expression induced a decrease in mRNA expression of *CIS*, *pim-1*, and *oncostatin M* (Figure 4F, lower panel). DN Stat5 suppressed in vitro proliferation of GFP-expressing dko KSL cells (designated as dko KSL/DNStat5 cells) more strongly than that of WT KSL cells (Figure 4G), demonstrating that dko KSL cells not only require Stat5 activity for their growth, but they are also more sensitive to Stat5 inhibition than WT cells. By contrast, DN Stat3 had little effect on the prolif-

eration of dko or WT KSL cells. Colony-forming assays showed a drastic reduction of myeloid colony (M-CFU and G-CFU) formation by dko KSL/DNStat5 cells, whereas dko KSL/DNStat3 cells exhibited a capacity to form various myeloid colonies similar to dko KSL/vector cells (Figure 4H). Next, the transduced dko KSL cells were transferred to irradiated C57BL/6-Ly5.1 mice. We monitored donor-derived cells in peripheral blood at 3 and 9 weeks. As loss of Stat5 was shown to result in reduced HSC engraftment (29), proportions of donor-derived cells in dko KSL/DNStat5 recipient mice were 3- to 4-fold lower than in dko KSL/vector recipient mice

**Figure 5**

Stat5 activation is required for the increased proliferation and macrophage differentiation of *SHIP*^{-/-} KSL cells. **(A)** KSL CD34⁺ cells derived from WT and *SHIP*^{-/-} mice were sorted into a 96-well plate containing SCF, Flt3 ligand, and IL-11 and cultured for 3 days. **P* < 0.05 between WT and mutant. **(B)** Colony-forming assays showed skewed differentiation of KSL cells of *SHIP*^{-/-} mice into macrophages in the presence of IL-3, IL-6, SCF, and erythropoietin (left panel). *SHIP*^{-/-} KSL cells formed colonies in the absence of growth factors (right panel). **(C–E)** Stat5 in *SHIP*^{-/-} c-Kit⁺Lin⁻ cells **(C)** and F4/80⁺ lung macrophages **(C–E)** was constitutively activated, as shown by flow cytometry **(C)**, immunohistochemistry **(D)**, and confocal microscopy **(E)** using anti-p-Stat5 (Tyr694). Original magnification, ×400 **(D and E)**. **(F and G)** *SHIP*^{-/-} and WT KSL cells were transduced with a bicistronic retroviral vector encoding DN Stat5, DN Stat3, or empty vector, together with GFP, and cultured in SCF, Flt3 ligand, and IL-11. Ratios of cell numbers normalized against those of empty vector-transduced cells are plotted. #*P* < 0.05 compared with cells transduced with empty vector. **(F)** Transduced cells (150 cells seeded) were cultured to form colonies in methylcellulose medium containing IL-3, IL-6, SCF, and erythropoietin **(G)**.

(10.0% ± 0.8% versus 33.0% ± 13% at 3 weeks; 9.0% ± 1.8% versus 38.0% ± 14% at 9 weeks; *P* < 0.05 at both time points). We also observed lower proliferation/differentiation of dko KSL/DNStat5 cells in the recipient mice, as 68.0% ± 8.7% of the donor-derived cells were CD11b⁺ (granulocytes and/or monocytes) in dko KSL/vector mice 9 weeks after transplantation, while only 21.4% ± 6.2% (*P* = 0.002) were CD11b⁺ in dko KSL/DNStat5 mice. Nine weeks after transplant, dko KSL/vector, but not dko KSL/DNStat5, mice developed MPD and lung inflammation (Table 1). Therefore, the lack of MPD in dko KSL/DNStat5 recipient mice can be due to the low engraftment or combined effects of low engraftment and low proliferation/differentiation of these cells. These results collectively demonstrate that Stat5 activation is required for the increased proliferation and myeloid differentiation activity of dko HSCs, leading to the development of MPD and lung disease.

MPD and lung phenotypes and their dependence on Stat5 are shared by SHIP- and Lyn/Hck-deficient mice. Comparison of our analysis of Lyn/Hck-deficient mice with previous studies on *SHIP*^{-/-} mice (15, 17, 30), some of whose properties were confirmed in this study (Supplemental Figure 8), suggests that there is a phenocopy between *lyn/hck* dko and *SHIP*^{-/-} mice in terms of MPD,

lung disease, and perturbed hematopoiesis. HSCs and GMPs were increased, but common lymphoid progenitors were reduced in BM of *SHIP*^{-/-} mice (Supplemental Figure 9A). *SHIP*^{-/-} HSCs also proliferated better than WT counterparts (Figure 5A), accompanied by accelerated cell cycling (Supplemental Figure 9B) in agreement with a recent study (12). *SHIP*^{-/-} HSCs formed more monocyte/macrophage colonies in the presence of cytokines than WT cells and also formed colonies spontaneously in the absence of growth factors (Figure 5B). The MPD and lung disease characterized by massive M2 macrophage infiltration were also transplantable with LT-HSCs from BM of *SHIP*^{-/-} mice (Supplemental Figure 9D). Stat5 phosphorylation was increased in both HSCs and BAL macrophages from *SHIP*^{-/-} mice (Figure 5, C–E). Consistent with this, mRNA expression of Stat5 target genes was upregulated in *SHIP*^{-/-} HSCs (Supplemental Figure 9C). Finally, introduction of DN Stat5 into *SHIP*^{-/-} HSCs inhibited the proliferation and monocyte/macrophage colony formation of KSL cells (Figure 5, F and G). DN Stat5 expression prevented the development of MPD and lung disease upon transplantation of *SHIP*^{-/-} KSL cells (Supplemental Figure 9D), demonstrating that the MPD and lung disease in *SHIP*^{-/-} mice are also Stat5-dependent.

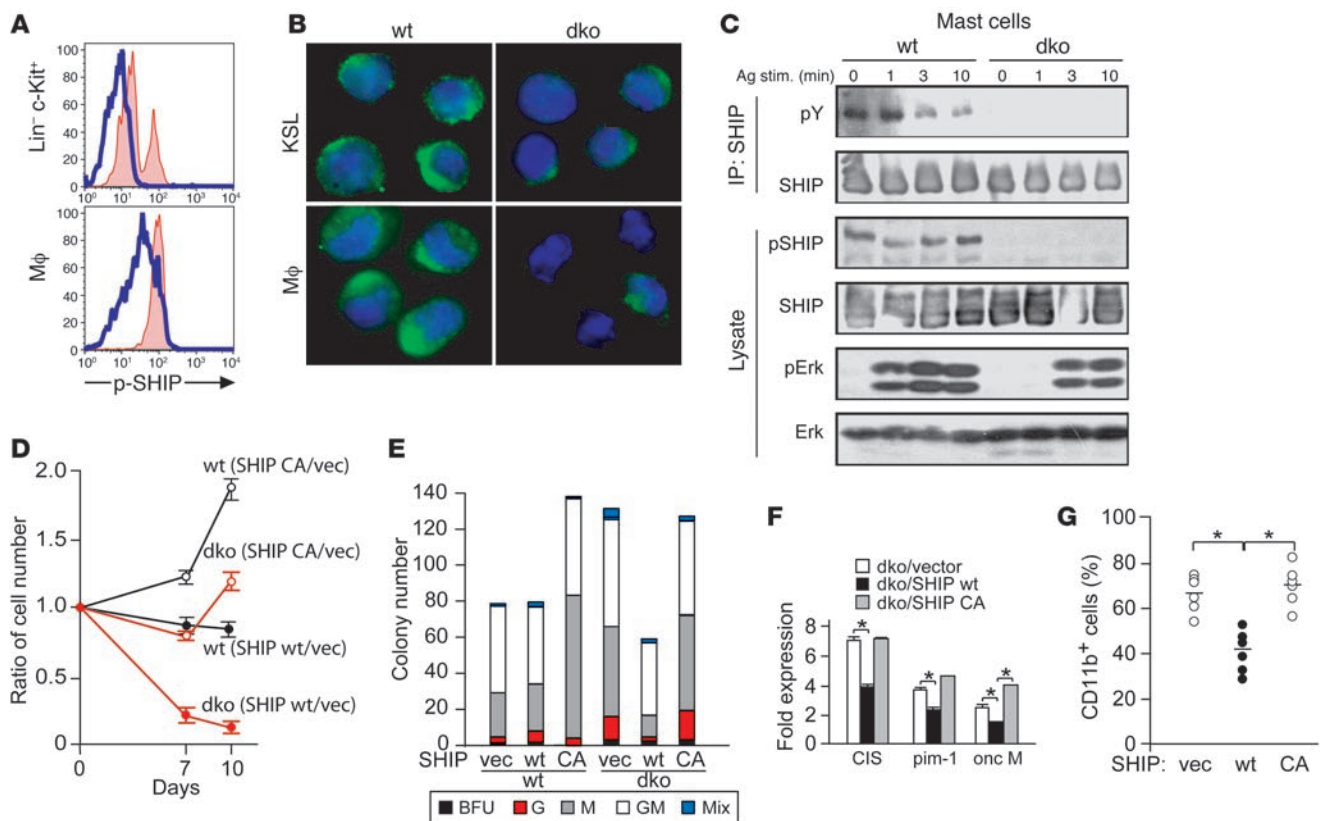


Figure 6

SHIP phosphorylation is reduced in *Lyn/Hck*-deficient HSCs and expression of a membrane-bound form of SHIP reverses dko phenotypes. (A and B) SHIP phosphorylation in KSL cells and macrophages was lower in dko than WT mice, as shown by flow cytometric analysis (A) and confocal microscopy (B) using anti-p-SHIP (Tyr1020). Original magnification, $\times 400$. (C) WT and dko BM-derived mast cells were sensitized with anti-DNP IgE and stimulated with antigen for the indicated periods. Tyrosine phosphorylation of SHIP was analyzed by immunoprecipitation followed by probing with anti-phosphotyrosine mAb (pY, upper panel) or cell lysates were directly analyzed by Western blotting using anti-p-SHIP (Tyr1020) antibody (lower panel). As a control to confirm activation of cells, blots were probed with anti-p-ERK1/2. (D and E) dko and WT KSL cells were transduced with a bicistronic retroviral vector encoding rCD2/SHIP WT, rCD2/SHIP C701A (CA), or empty vector, together with GFP, and cultured in medium containing SCF, Flt3 ligand, and IL-11. Ratios of cell numbers normalized against those of empty vector-transduced cells are plotted (D). Three days after retrovirus infection, some transduced cells (150 cells seeded) were cultured in methylcellulose containing IL-3, IL-6, SCF, and erythropoietin (E). (F) Real-time PCR analysis of mRNA expression of Stat5 target genes in dko KSL cells transduced with rCD2/SHIP WT or empty vector. (G) Dko KSL cells transduced with a bicistronic retroviral vector encoding rCD2/SHIP WT, rCD2/SHIP CA, or empty vector were i.v. injected into lethally irradiated mice. Eight weeks later, CD11b⁺GFP⁺ cells in peripheral blood were enumerated by flow cytometry. * $P < 0.05$.

*Expression of membrane-bound SHIP ameliorates or prevents MPD and lung disease in *Lyn/Hck*-deficient mice.* The similarities between SHIP^{-/-} and *Lyn/Hck*-deficient mice raised the possibility that SHIP may be a mediator linking *Lyn/Hck* to activated Stat5. Indeed, we observed little SHIP phosphorylation in KSL, BAL macrophages, and BM-derived mast cells from dko mice (Figure 6, A–C), indicating that SHIP is a downstream target of *Lyn* and *Hck*. However, SHIP phosphorylation levels were comparable in *lyn*^{-/-}, *hck*^{-/-}, and WT mice (data not shown), suggesting that *Lyn* and *Hck* can compensate for each other and phosphorylate SHIP. The recruitment of SHIP to the plasma membrane is phosphorylation dependent and is a prerequisite for its function (31). Since SHIP phosphorylation and thus its membrane recruitment did not take place in dko cells, we reasoned that expression of a membrane-bound form of SHIP might reverse the phenotype of dko mice. To test this possibility, we took advantage of expression constructs encoding a rat CD2/SHIP fusion protein, in which the intracellular por-

tion of rat CD2 was replaced by the SHIP phosphatase core of either WT or catalytically inactive (C701A) mutant enzyme. The fusion proteins were previously shown to constitutively reside in the plasma membrane (32). Expression of rCD2/SHIP WT in dko KSL cells inhibited their proliferative and colony (M-CFU and G-CFU) forming abilities (Figure 6, D and E). The development of lung disease was spared in recipient mice of rCD2/SHIP WT-transduced dko KSL cells, although CD11b⁺ cells in their peripheral blood were reduced but still higher than in WT mice (Figure 6G and data not shown). By contrast, rCD2/SHIP C701A did not inhibit the proliferative or colony-forming activities of dko HSCs or in vivo disease development. Interestingly, expression of rCD2/SHIP C701A rather increased the proliferative and monocyte/macrophage colony-forming abilities of WT HSCs (Figure 6, D and E), indicating that rCD2/SHIP C701A has a DN function. These DN effects were not observed in dko cells, probably because SHIP functions in dko cells were already very low.

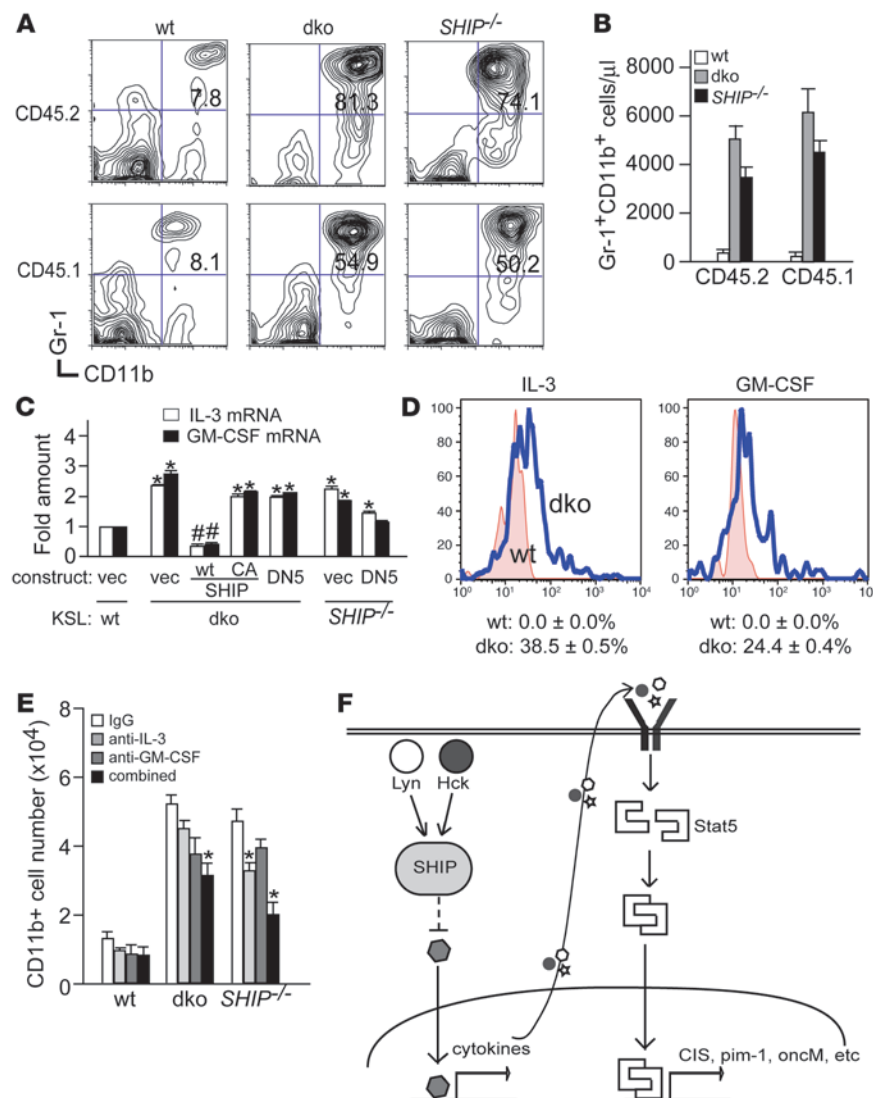


Figure 7

Cytokines secreted from dko and *SHIP*^{-/-} cells promote in vitro and in vivo expansion of myeloid cells in an autocrine/paracrine manner. (A and B) In vivo paracrine effect of dko and *SHIP*^{-/-} cells. Equal numbers of CD45.1⁺ (Ly5.1) BM cells and CD45.2⁺ (Ly5.2) BM cells derived from either WT, dko, or *SHIP*^{-/-} mice were cotransferred into lethally irradiated Ly5.1 recipients. Eight weeks later, CD45.1 or CD45.2 cells in peripheral blood were gated and Gr-1⁺CD11b⁺ myeloid cells were enumerated. Percentages of Gr-1⁺CD11b⁺ cells within each CD45 cohort are indicated. (C) RNAs from WT, dko, or *SHIP*^{-/-} KSL cells transduced with the indicated bicistronic vectors were submitted to real-time PCR analysis. **P* < 0.05 versus WT/vector, #*P* < 0.05 versus dko/vector values, by Student's *t* test. (D) Intracellular staining of IL-3 and GM-CSF in WT and dko KSL cells. (E) Lin⁻ cells (Ly5.1) were cocultured for a week with either WT, dko, or *SHIP*^{-/-} Lin⁻ cells (Ly5.2⁺) in the presence of control IgG, anti-IL-3, anti-GM-CSF, or anti-IL-3 plus anti-GM-CSF. CD11b⁺ cells were enumerated by flow cytometry. (F) A proposed model. Both Lyn and Hck are required to fully activate SHIP that in turn inhibits the production of several cytokines. In the absence of Lyn and Hck, secreted cytokines, including IL-3 and GM-CSF, can activate the Jak2/Stat5 pathway, eventually disrupting the homeostasis of HSCs.

Furthermore, rCD2/SHIP WT expression downregulated mRNA expression of Stat5 target genes (Figure 6F), suggesting that the activity of Stat5 in dko HSCs was at least partially inhibited by rCD2/SHIP WT. Therefore, we concluded that SHIP is a mediator linking Lyn/Hck to Stat5.

IL-3 and GM-CSF secreted from Lyn/Hck- and SHIP-deficient HSCs/progenitors promote the expansion of myeloid cells. The above results suggest that Lyn and Hck activate SHIP and that SHIP in turn inhibits Stat5 activity. Previous studies showed that SHIP negatively regulates the production of several cytokines (15, 33). Therefore, it is possible that SHIP either directly inhibits Stat5 activity or indirectly inhibits Stat5 activity by inhibiting the secretion of a cytokine(s) that induces the activation of Stat5 by an autocrine or paracrine mechanism. Direct SHIP-dependent regulation of Stat5 activity seems unlikely, as expression of rCD2/SHIP WT or C701A did not affect IL-3-induced Stat5 phosphorylation in Ba/F3 mouse pro-B cells (data not shown). Consistent with the paracrine mechanism, cotransfer of dko or *SHIP*^{-/-} BM cells (Ly5.2) together with WT BM cells (Ly5.1) into irradiated Ly5.1 mice revealed that dko and *SHIP*^{-/-} cells can promote the expansion of myeloid cells that differentiated from cotransferred Ly5.1⁺ WT

cells (Figure 7, A and B). We also detected increased mRNA expression of several cytokines including IL-3, GM-CSF and G-CSF in dko KSL cells (data not shown). Importantly, transduction with rCD2/SHIP WT, but not rCD2/SHIP C701A, abrogated the expression of these cytokines in dko KSL cells (Figure 7C). By contrast, transduction with DN Stat5 only slightly reduced the expression of these genes in dko and *SHIP*^{-/-} KSL cells, indicating that the increased expression of these cytokine genes is largely independent of Stat5 activity. Intracellular staining showed that KSL and lineage⁻, but not lineage⁺, cells of dko mice express IL-3 and GM-CSF (Figure 7D and data not shown). IL-3 and GM-CSF have been known to activate Stat5 (34). Incubation with IL-3 or GM-CSF promotes myeloid differentiation and proliferation of lineage⁻ cells in our culture conditions (data not shown). To test whether IL-3 and GM-CSF secreted from dko or *SHIP*^{-/-} cells are involved in the myeloid expansion of cotransferred WT Ly5.1 cells, we cocultured lineage⁻ Ly5.1 cells with either WT, dko or *SHIP*^{-/-} cells (Ly5.2) in vitro. One week later, CD11b⁺ cells in the Ly5.1 pool were increased by approximately 3-fold when cocultured with dko and *SHIP*^{-/-} cells compared with those cocultured with Ly5.2⁺ WT cells (Figure 7E). Addition of anti-IL-3 or anti-GM-



CSF neutralizing antibodies alone slightly reduced the numbers of CD11b⁺ cells in the Ly5.1 pool, but the inhibitory effect of the combined antibodies was more substantial, reaching 40%–50% inhibition (Figure 7E). This inhibition could be larger, as control experiments showed that the antibodies we used blocked IL-3- and GM-CSF-induced expansion of CD11b⁺ cells at efficiencies of 76% and 72%, respectively. These results demonstrated that paracrine secretion of IL-3 and GM-CSF is largely responsible for the expansion of myeloid cells of Ly5.1⁺ WT in cocultures with dko and *SHIP*^{-/-} cells. Furthermore, anti-IL-3 and anti-GM-CSF antibodies also inhibited growth factor-independent myeloid colony formation from dko BM cells (Figure 3F).

Discussion

In this study we revealed a novel signaling pathway that when perturbed can lead to myeloproliferative transformation: Lyn and Hck activate SHIP; active SHIP in turn inhibits autocrine or paracrine production of cytokines including IL-3 and GM-CSF, which, without upstream control, would lead to chronically activated Stat5 signaling (Figure 7F). This pathway is essential for the homeostasis of HSCs. We demonstrated that deficiency in Lyn/Hck or SHIP results in increased proliferation, growth factor-independent growth, and myeloid-skewed differentiation of HSCs, eventually generating transformed myeloid cells. Both the MPD and lung disease in Lyn/Hck- and SHIP-deficient mice seem to originate from the LT-HSC compartment.

SFKs can be activated by various cytokines and growth factors including IL-3 and GM-CSF. Previous studies support the sensitivity of this pathway to myeloproliferative transformation: GM-CSF-transgenic mice develop an extensive accumulation of macrophages in various organs (35). Transgenic mice expressing constitutively active mutants of the common β subunit of the receptors for IL-3, IL-5, and GM-CSF induce MPD (36, 37). Interestingly, adoptive transfer of Nf1 tumor suppressor-deficient fetal liver cells induces MPD, a mouse model of human juvenile myelomonocytic leukemia, and GM-CSF and the common β receptor subunit are involved in establishing and maintaining this MPD (38, 39). Constitutively active Stat5 in KSL CD34⁺ cells can induce a fatal MPD (40). Several oncogenes, including BCR-ABL, FLT3 ITD, and JAK2 V617F, can induce MPD or frank leukemia in part by activating Stat5 at HSC levels (41–46). Interestingly, both the growth and monocyte/macrophage colony-forming activities in dko and *SHIP*^{-/-} HSCs depend more heavily on Stat5 than in WT HSCs. This phenomenon may represent a case of “oncogene addiction” of the mutant cells (47).

As described in the Introduction, Lyn and Hck play negative regulatory roles in myeloid cell differentiation in mice. Unlike *lyn/hck* dko mice, *lyn/hck/fgr* triple knockout mice did not develop MPD or dramatic abnormalities in hematopoiesis (48). This might suggest that Fgr antagonizes our proposed pathway (Figure 7), although it is not known at what level Fgr interferes with the pathway. However, it is possible that Fgr antagonizes Lyn and Hck by an unknown mechanism. As these SFKs, SHIP, cytokines, and Stat5 are all conserved in humans, our proposed pathway is likely operative in human hematopoietic cells. However, gaps in our knowledge of SFKs' function in hematopoiesis are illustrated in their pathogenetic role in human chronic myelogenous leukemia (CML): both human LYN and HCK could phosphorylate BCR-ABL and contribute to CML (49–51). Furthermore, imatinib-resistant BCR-ABL⁺ K562 cells expressed a higher level of LYN,

and knocking down of LYN could reverse this resistance (52, 53). In a mouse model, however, Lyn, Hck, and Fgr are required for BCR-ABL-induced B cell acute lymphoblastic leukemia, but not BCR-ABL-induced CML (54). By contrast, our data demonstrated that Lyn and Hck are important negative regulators for a CML-like disease (e.g., MPD) in mice, raising such questions as whether in vivo functions of LYN and HCK in human CML are largely different from their in vitro functions or whether the roles of Lyn and Hck are species-specific. These issues are important to address in light of recent applications of SFK inhibitors in the treatment of imatinib-resistant or -intolerant CML (55, 56).

Similar to SHIP, phosphorylation of SHP-1 is also Lyn-dependent (57, 58), and it was totally lost in dko cells (data not shown). SHP-1 can interact with, and inhibit phosphorylation of, Stat5 (59–61). Importantly, motheaten mice with homozygous mutations in the *SHP-1* gene have a phenotype similar to dko and *SHIP*^{-/-} mice, in terms of MPD and lung pathology (62, 63). It will be interesting to investigate whether SHP-1 plays any role in the pathogenesis of the MPD and lung disease in dko mice.

Lung inflammation appears to be a cause of premature deaths in dko and *SHIP*^{-/-} mice. The lung disease is characterized by a dramatic increase in M2 macrophages within alveoli. M2 macrophages are involved in debris scavenging, angiogenesis, tissue remodeling, wound healing, and the promotion of type II immunity, whereas classically activated (M1) macrophages eradicate invading microorganisms and tumor cells and promote type I immune responses (13, 14). Thus, M2 macrophages in dko and *SHIP*^{-/-} lungs likely contribute to the pathogenesis of lung disease. However, little is known about the mechanisms underlying macrophage programming, and it is unresolved whether M1 and M2 macrophages represent distinct, defined lineages or whether mature macrophage phenotypes are plastic and adaptable to ever-changing environments. Our data confirmed and extended a previous study (15) by indicating that Lyn/Hck and SHIP are important for M2 macrophage programming. Increased Stat5 activity may also play a role in promoting M2 macrophage differentiation, since constitutively active Stat5 could induce Ba/F3 cells to differentiate into macrophages (64), and Stat5 was reported to be required for FDC-P1 cell differentiation into monocytes (65).

Methods

Mice. *Lyn*^{-/-} (3), *hck*^{-/-} (66), and *SHIP*^{-/-} (10) mice were described previously. Dko mice were bred from *lyn*^{-/-} and *hck*^{-/-} mice. These mice were backcrossed with C57BL/6 mice for 15 or more generations. The CD45 congenic mouse strains C57BL/6-Ly5.1 mice were purchased from The Jackson laboratory. Animal studies were approved by the institutional review board of La Jolla Institute for Allergy and Immunology.

Identification and purification of HSCs and progenitors. Cell sorting was performed by immunomagnetic-based pre-enrichment followed by multicolor flow cytometric sorting (see Supplemental Data for details).

In vitro cultures and colony-forming assays. LT-HSCs or Lin⁻ cells were sorted into a 96-well round bottom plate (1 cell per well) in 200 μ l of IMDM containing 5% FBS, 50 μ M 2-mercaptoethanol, and a cocktail of SCF, IL-11, and Flt3 ligand. For myeloid colony-forming assays, BM cells or HSCs were cultured in Methocult M3434 (StemCell Technologies). For lymphoid colony-forming assays, BM cells were cultured in Methocult M3630 (StemCell Technologies). After 7–12 days, colonies were scored by microscopy, Wright-Giemsa staining, and flow cytometric analysis.

Transplantation of hematopoietic cells. BM cells were injected into lethally (9.6 Gy) irradiated recipient mice. For transplantation of HSCs and pro-



genitors, sorted LT-HSCs, ST-HSCs, CMPs, GMPs, and retrovirally transduced HSCs were injected into lethally irradiated mice together with Sca-1-depleted C57BL/6-Ly5.1 helper BM cells.

Statistics. Two-tailed Student's *t* test was used throughout this study. A *P* value of less than 0.05 was considered significant.

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